RAW SEQUENCE LISTING

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Application Serial Number: _10/579,286_

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RAW SEQUENCE LISTING   DATE: 05/24/2006
PATENT APPLICATION: **US/10/579,286** TIME: 12:21:38

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Output Set: **N:\CRF4\05242006\J579286.raw**

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     3 <110> APPLICANT: ANDREEV, et al.
     5 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING MODULATORS OF ACTIVE KIT
TYROSINE KINASE
     6         RECEPTOR
     8 <130> FILE REFERENCE: 30694/39618A
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/579,286
C--> 10 <141> CURRENT FILING DATE: 2006-05-15
    10 <150> PRIOR APPLICATION NUMBER: US 60/526,930
    11 <151> PRIOR FILING DATE: 2003-12-04
    13 <160> NUMBER OF SEQ ID NOS: 6
    15 <170> SOFTWARE: PatentIn version 3.2
    17 <210> SEQ ID NO: 1
    18 <211> LENGTH: 5084
    19 <212> TYPE: DNA
    20 <213> ORGANISM: Homo sapiens
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    24 <222> LOCATION: (22)..(2952)
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    31 ctc tgc gtt ctg ctc cta ctg ctt cgc gtc cag aca ggc tct tct caa       99
    32 Leu Cys Val Leu Leu Leu Leu Leu Arg Val Gln Thr Gly Ser Ser Gln
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    35 cca tct gtg agt cca ggg gaa ccg tct cca cca tcc atc cat cca gga      147
    36 Pro Ser Val Ser Pro Gly Glu Pro Ser Pro Pro Ser Ile His Pro Gly
    37             30                  35                  40
    39 aaa tca gac tta ata gtc cgc gtg ggc gac gag att agg ctg tta tgc      195
    40 Lys Ser Asp Leu Ile Val Arg Val Gly Asp Glu Ile Arg Leu Leu Cys
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    43 act gat ccg ggc ttt gtc aaa tgg act ttt gag atc ctg gat gaa acg      243
    44 Thr Asp Pro Gly Phe Val Lys Trp Thr Phe Glu Ile Leu Asp Glu Thr
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    47 aat gag aat aag cag aat gaa tgg atc acg gaa aag gca gaa gcc acc      291
    48 Asn Glu Asn Lys Gln Asn Glu Trp Ile Thr Glu Lys Ala Glu Ala Thr
    49 75                  80                  85                  90
    51 aac acc ggc aaa tac acg tgc acc aac aaa cac ggc tta agc aat tcc      339
    52 Asn Thr Gly Lys Tyr Thr Cys Thr Asn Lys His Gly Leu Ser Asn Ser
    53                 95                  100                 105
    55 att tat gtg ttt gtt aga gat cct gcc aag ctt ttc ctt gtt gac cgc      387
    56 Ile Tyr Val Phe Val Arg Asp Pro Ala Lys Leu Phe Leu Val Asp Arg
    57                 110                 115                 120
    59 tcc ttg tat ggg aaa gaa gac aac gac acg ctg gtc cgc tgt cct ctc      435
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63  aca gac cca gaa gtg acc aat tat tcc ctc aag ggg tgc cag ggg aag    483
64  Thr Asp Pro Glu Val Thr Asn Tyr Ser Leu Lys Gly Cys Gln Gly Lys
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67  cct ctt ccc aag gac ttg agg ttt att cct gac ccc aag gcg ggc atc    531
68  Pro Leu Pro Lys Asp Leu Arg Phe Ile Pro Asp Pro Lys Ala Gly Ile
69  155                 160                 165                 170
71  atg atc aaa agt gtg aaa cgc gcc tac cat cgg ctc tgt ctg cat tgt    579
72  Met Ile Lys Ser Val Lys Arg Ala Tyr His Arg Leu Cys Leu His Cys
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79  aaa gtg agg cca gcc ttc aaa gct gtg cct gtt gtg tct gtg tcc aaa    675
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83  gca agc tat ctt ctt agg gaa ggg gaa gaa ttc aca gtg acg tgc aca    723
84  Ala Ser Tyr Leu Leu Arg Glu Gly Glu Glu Phe Thr Val Thr Cys Thr
85          220                 225                 230
87  ata aaa gat gtg tct agt tct gtg tac tca acg tgg aaa aga gaa aac    771
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91  agt cag act aaa cta cag gag aaa tat aat agc tgg cat cac ggt gac    819
92  Ser Gln Thr Lys Leu Gln Glu Lys Tyr Asn Ser Trp His His Gly Asp
93                  255                 260                 265
95  ttc aat tat gaa cgt cag gca acg ttg act atc agt tca gcg aga gtt    867
96  Phe Asn Tyr Glu Arg Gln Ala Thr Leu Thr Ile Ser Ser Ala Arg Val
97                  270                 275                 280
99  aat gat tct gga gtg ttc atg tgt tat gcc aat aat act ttt gga tca    915
100 Asn Asp Ser Gly Val Phe Met Cys Tyr Ala Asn Asn Thr Phe Gly Ser
101             285                 290                 295
103 gca aat gtc aca aca acc ttg gaa gta gta gat aaa gga ttc att aat    963
104 Ala Asn Val Thr Thr Thr Leu Glu Val Val Asp Lys Gly Phe Ile Asn
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107 atc ttc ccc atg ata aac act aca gta ttt gta aac gat gga gaa aat   1011
108 Ile Phe Pro Met Ile Asn Thr Thr Val Phe Val Asn Asp Gly Glu Asn
109 315                 320                 325                 330
111 gta gat ttg att gtt gaa tat gaa gca ttc ccc aaa cct gaa cac cag   1059
112 Val Asp Leu Ile Val Glu Tyr Glu Ala Phe Pro Lys Pro Glu His Gln
113                 335                 340                 345
115 cag tgg atc tat atg aac aga acc ttc act gat aaa tgg gaa gat tat   1107
116 Gln Trp Ile Tyr Met Asn Arg Thr Phe Thr Asp Lys Trp Glu Asp Tyr
117             350                 355                 360
119 ccc aag tct gag aat gaa agt aat atc aga tac gta agt gaa ctt cat   1155
120 Pro Lys Ser Glu Asn Glu Ser Asn Ile Arg Tyr Val Ser Glu Leu His
121             365                 370                 375
123 cta acg aga tta aaa ggc acc gaa gga ggc act tac aca ttc cta gtg   1203
124 Leu Thr Arg Leu Lys Gly Thr Glu Gly Gly Thr Tyr Thr Phe Leu Val
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131 aca aaa cca gaa atc ctg act tac gac agg ctc gtg aat ggc atg ctc    1299
132 Thr Lys Pro Glu Ile Leu Thr Tyr Asp Arg Leu Val Asn Gly Met Leu
133                    415                   420                   425
135 caa tgt gtg gca gca gga ttc cca gag ccc aca ata gat tgg tat ttt    1347
136 Gln Cys Val Ala Ala Gly Phe Pro Glu Pro Thr Ile Asp Trp Tyr Phe
137                    430                   435                   440
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140 Cys Pro Gly Thr Glu Gln Arg Cys Ser Ala Ser Val Leu Pro Val Asp
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143 gtg cag aca cta aac tca tct ggg cca ccg ttt gga aag cta gtg gtt    1443
144 Val Gln Thr Leu Asn Ser Ser Gly Pro Pro Phe Gly Lys Leu Val Val
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147 cag agt tct ata gat tct agt gca ttc aag cac aat ggc acg gtt gaa    1491
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149 475                    480                   485                   490
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156 Ala Phe Lys Gly Asn Asn Lys Glu Gln Ile His Pro His Thr Leu Phe
157                    510                   515                   520
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161              525                   530                   535
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164 Ile Val Met Ile Leu Thr Tyr Lys Tyr Leu Gln Lys Pro Met Tyr Glu
165      540                   545                   550
167 gta cag tgg aag gtt gtt gag gag ata aat gga aac aat tat gtt tac    1731
168 Val Gln Trp Lys Val Val Glu Glu Ile Asn Gly Asn Asn Tyr Val Tyr
169 555                    560                   565                   570
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175 aac agg ctg agt ttt ggg aaa acc ctg ggt gct gga gct ttc ggg aag    1827
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177                    590                   595                   600
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180 Val Val Glu Ala Thr Ala Tyr Gly Leu Ile Lys Ser Asp Ala Ala Met
181              605                   610                   615
183 act gtc gct gta aag atg ctc aag ccg agt gcc cat ttg aca gaa cgg    1923
184 Thr Val Ala Val Lys Met Leu Lys Pro Ser Ala His Leu Thr Glu Arg
185      620                   625                   630
187 gaa gcc ctc atg tct gaa ctc aaa gtc ctg agt tac ctt ggt aat cac    1971
188 Glu Ala Leu Met Ser Glu Leu Lys Val Leu Ser Tyr Leu Gly Asn His
189 635                    640                   645                   650
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195 ctg gtc att aca gaa tat tgt tgc tat ggt gat ctt ttg aat ttt ttg    2067
196 Leu Val Ile Thr Glu Tyr Cys Cys Tyr Gly Asp Leu Leu Asn Phe Leu
197             670                 675                 680
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203 gaa gct gca ctt tat aag aat ctt ctg cat tca aag gag tct tcc tgc    2163
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212 Val Val Pro Thr Lys Ala Asp Lys Arg Arg Ser Val Arg Ile Gly Ser
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215 tac ata gaa aga gat gtg act ccc gcc atc atg gag gat gac gag ttg    2307
216 Tyr Ile Glu Arg Asp Val Thr Pro Ala Ile Met Glu Asp Asp Glu Leu
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219 gcc cta gac tta gaa gac ttg ctg agc ttt tct tac cag gtg gca aag    2355
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228 Ala Arg Asn Ile Leu Leu Thr His Gly Arg Ile Thr Lys Ile Cys Asp
229 795                 800                 805                 810
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232 Phe Gly Leu Ala Arg Asp Ile Lys Asn Asp Ser Asn Tyr Val Val Lys
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235 gga aac gct cga cta cct gtg aag tgg atg gca cct gaa agc att ttc    2547
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248 Val Asp Ser Lys Phe Tyr Lys Met Ile Lys Glu Gly Phe Arg Met Leu
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251 agc cct gaa cac gca cct gct gaa atg tat gac ata atg aag act tgc    2739
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272 Leu Val His Asp Asp Val
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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date